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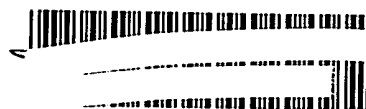
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430

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p#3

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/082,659

DATE: 03/13/2002

TIME: 11:05:58

Input Set : N:\Crf3\RULE60\10082659.raw

Output Set: N:\CRF3\03132002\J082659.raw

ENTERED

RECEIVED

APR 11 2002

TECH CENTER 1600/2900

```

1 <110> APPLICANT: Abbott Laboratories
2   Cohen, Maurice
3   Colpitts, Tracey L.
4   Friedman, Paula N.
5   Granados, Edward N.
6   Klass, Michael R.
7   Russell, John C.
8   Stroupe, Stephen D.
9 <120> TITLE OF INVENTION: REAGENTS AND METHODS USEFUL FOR
10  DETECTING DISEASES OF THE PROSTATE
11 <130> FILE REFERENCE: 6171.US.P1
13 <140> CURRENT APPLICATION NUMBER: 10/082,659
14 <141> CURRENT FILING DATE: 2002-02-25
16 <150> PRIOR APPLICATION NUMBER: 09/566,876
17 <151> PRIOR FILING DATE: 2000-05-08
18 <160> NUMBER OF SEQ ID NOS: 21
19 <170> SOFTWARE: FastSEQ for Windows Version 3.0
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 187
23 <212> TYPE: DNA
24 <213> ORGANISM: Homo sapiens
25 <220> FEATURE:
26 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
27   position 2
28 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
29   position 96
30 <400> SEQUENCE: 1
W--> 31   tnttttttgt ttgctctgaa tttattgcga gtgaaaaaca gagaaaatcc tcaagtttaa      60
W--> 32   gtttctgata gcagagtgtg ggagtttagag catggngagt ccagagggtc cagaccccca      120
33   aaggctctcta ccagggccat ctccgttagt ggcggtggca gcccctcttg tggccttttt      180
34   cctctct
36 <210> SEQ ID NO: 2
37 <211> LENGTH: 250
38 <212> TYPE: DNA
39 <213> ORGANISM: Homo sapiens
40 <400> SEQUENCE: 2
41   gtttgctctg aatttattgc gagtgaaaaa cagagaaaat cctcaagttt aagtttctga      60
42   tagcagagtg tgggagttag agcatgggga gtccagaggt tccagacccc caaagggtctc      120
43   taccagggcc atctccgtta gtggcggttg cagcccctct tgtggccttt ttctctcttc      180
44   caagggggtca ccccgcacca tgccgctccc cctcatctat cttgccccct cccacagggc      240
45   ccatctgcgc
47 <210> SEQ ID NO: 3
48 <211> LENGTH: 224

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RAW.SEQUENCE LISTING

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```

49 <212> TYPE: DNA
50 <213> ORGANISM: Homo sapiens
51 <220> FEATURE:
52 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
53 position 71
54 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
55 position 73
56 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
57 position 76
58 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
59 position 81
60 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
61 position 93
62 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
63 position 138
64 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
65 position 187
66 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
67 positions 191 and 192
68 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
69 position 207
70 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
71 position 212
72 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
73 position 223
74 <400> SEQUENCE: 3
75      cgatcgggca agtaaaccce ctccctcgcc gacttcggaa ctggcgagag ttcagcgcag      60
W--> 76      atgggcctgt ngngangggg naagatagat ganggggagc ggcattgtgc ggggtgaccc      120
W--> 77      cttggagaga ggaaaaangc cacaagaggg gctgccaccg ccactaacgg agatggccct      180
W--> 78      ggtaganact nnggggggtct ggaactnctg gnetccccat gcnc      224
80 <210> SEQ ID NO: 4
81 <211> LENGTH: 312
82 <212> TYPE: DNA
83 <213> ORGANISM: Homo sapiens
84 <220> FEATURE:
85 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
86 position 2
87 <400> SEQUENCE: 4
W--> 88      tnttttttgt ttgctctgaa tttattgcga gtgaaaaaca gagaaaatcc tcaagtttaa      60
89      gtttctgata gcagagtgtg ggagtttagag catggggagt ccagaggttc cagacccccca      120
90      aaggtctcta ccagggccat ctccgttagt ggcggtggca gcccctcttg tgccctttt      180
91      cctctctcca aggggtcacc ccgcaccatg ccgtccccc tcactatct tgccccctcc      240
92      ccacaggccc atctgcgctg aactctcgcc agttccgaag tcggcgaggg aggggggtta      300
93      cttgcccgat cg      312
95 <210> SEQ ID NO: 5
96 <211> LENGTH: 68
97 <212> TYPE: DNA
98 <213> ORGANISM: Homo sapiens
99 <400> SEQUENCE: 5

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RAW SEQUENCE LISTING

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Input Set : N:\Crif3\RULE60\10082659.raw

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```

100      agctcggaat tccgagcttg gatacctctag agcggccgcc gactagttag ctcgtcgacc      60
101      cggaatt                                     68
103 <210> SEQ ID NO: 6
104 <211> LENGTH: 68
105 <212> TYPE: DNA
106 <213> ORGANISM: Homo sapiens
107 <400> SEQUENCE: 6
108      aattaattcc cgggtcgacg agctcactag tcggcggccg ctctagagga tccaagctcg      60
109      gaattccg                                     68
111 <210> SEQ ID NO: 7
112 <211> LENGTH: 24
113 <212> TYPE: DNA
114 <213> ORGANISM: Homo sapiens
115 <400> SEQUENCE: 7
116      agcggataac aatttcacac agga                                     24
118 <210> SEQ ID NO: 8
119 <211> LENGTH: 18
120 <212> TYPE: DNA
121 <213> ORGANISM: Homo sapiens
122 <400> SEQUENCE: 8
123      tgtaaaacga cggccagt                                     18
125 <210> SEQ ID NO: 9
126 <211> LENGTH: 73
127 <212> TYPE: PRT
128 <213> ORGANISM: Homo sapiens
129 <400> SEQUENCE: 9
130      Met Gly Ser Pro Glu Val Pro Asp Pro Gln Arg Ser Leu Pro Gly Pro
131      1          5          10          15
132      Ser Pro Leu Val Ala Val Ala Ala Pro Leu Val Ala Phe Phe Leu Ser
133      20          25          30
134      Pro Arg Gly His Pro Ala Pro Cys Arg Ser Pro Ser Ser Ile Leu Pro
135      35          40          45
136      Pro Pro His Arg Pro Ile Cys Ala Glu Leu Ser Pro Val Pro Lys Ser
137      50          55          60
138      Ala Arg Glu Gly Val Tyr Leu Pro Asp
139      65          70
141 <210> SEQ ID NO: 10
142 <211> LENGTH: 24
143 <212> TYPE: PRT
144 <213> ORGANISM: Homo sapiens
145 <400> SEQUENCE: 10
146      Met Gly Ser Pro Glu Val Pro Asp Pro Gln Arg Ser Leu Pro Gly Pro
147      1          5          10          15
148      Ser Pro Leu Val Ala Val Ala Ala
149      20
151 <210> SEQ ID NO: 11
152 <211> LENGTH: 31
153 <212> TYPE: PRT
154 <213> ORGANISM: Homo sapiens

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RAW SEQUENCE LISTING

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Input Set : N:\Crf3\RULE60\10082659.raw

Output Set: N:\CRF3\03132002\J082659.raw

```

155 <400> SEQUENCE: 11
156     Leu Val Ala Phe Phe Leu Ser Pro Arg Gly His Pro Ala Pro Cys Arg
157         1             5             10             15
158     Ser Pro Ser Ser Ile Leu Pro Pro Pro His Arg Pro Ile Cys Ala
159         20             25             30
161 <210> SEQ ID NO: 12
162 <211> LENGTH: 21
163 <212> TYPE: PRT
164 <213> ORGANISM: Homo sapiens
165 <400> SEQUENCE: 12
166     Ser Pro Arg Gly His Pro Ala Pro Cys Arg Ser Pro Ser Ser Ile Leu
167         1             5             10             15
168     Pro Pro Pro His Arg
169         20
171 <210> SEQ ID NO: 13
172 <211> LENGTH: 29
173 <212> TYPE: PRT
174 <213> ORGANISM: Homo sapiens
175 <400> SEQUENCE: 13
176     Ser Ile Leu Pro Pro Pro His Arg Pro Ile Cys Ala Glu Leu Ser Pro
177         1             5             10             15
178     Val Pro Lys Ser Ala Arg Glu Gly Val Tyr Leu Pro Asp
179         20             25
181 <210> SEQ ID NO: 14
182 <211> LENGTH: 13
183 <212> TYPE: PRT
184 <213> ORGANISM: Homo sapiens
185 <400> SEQUENCE: 14
186     Val Pro Lys Ser Ala Arg Glu Gly Val Tyr Leu Pro Asp
187         1             5             10
189 <210> SEQ ID NO: 15
190 <211> LENGTH: 8
191 <212> TYPE: PRT
192 <213> ORGANISM: Homo sapiens
193 <400> SEQUENCE: 15
194     Asp Tyr Lys Asp Asp Asp Asp Lys
195         1             5
197 <210> SEQ ID NO: 16
198 <211> LENGTH: 21
199 <212> TYPE: PRT
200 <213> ORGANISM: Homo sapiens
201 <400> SEQUENCE: 16
202     Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Met His Thr Glu His
203         1             5             10             15
204     His His His His His
205         20
207 <210> SEQ ID NO: 17
208 <211> LENGTH: 246
209 <212> TYPE: DNA

```

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```

210 <213> ORGANISM: Homo sapiens
211 <400> SEQUENCE: 17
212      tttttgtttg ctctgaattt attgcgagtg aaaaacagag aaaatcctca agtttaagtt      60
213      tctgatagca gagtgtggga gtagagcatg gggagtccag aggttccaga ccccccagg      120
214      tctctaccag gggcatctcc gttagtggcg gtggcagccc ctcttggtgc ctttttctc      180
215      tctccaagg gtcacccgc accatgccgc tccccctcat ctatcttgcc ccctccccac      240
216      aggcca      246
218 <210> SEQ ID NO: 18
219 <211> LENGTH: 704
220 <212> TYPE: DNA
221 <213> ORGANISM: Homo sapiens
222 <400> SEQUENCE: 18
223      tttttgtttg ctctgaattt attgcgagtg aaaaacagag aaaatcctca agtttaagtt      60
224      tctgatagca gagtgtggga gttagagcat gggagtcca gaggttccag acccccagg      120
225      gtctctacca gggccatctc cgttagtggc ggtggcagcc cctcttggtg ctttttctc      180
226      ctctccaagg ggtcacccgc caccatgccg ctccccctca tctatcttgc ccctccccca      240
227      caggcccatc tgcgtgaac tctcgccagt tccgaagtcg gcgaggagg gggtttactt      300
228      gccgatcgt tgggtgggtt gagcttatag aggagagga gtaagaacct gcgatattga      360
229      aagctacca catggggctt ccttgaagga ggacgtggaa ggcagaaagt gacctgctct      420
230      gagcggcgca tgtaaccgag gaccttaagc tggaccacgg ggcttggacg attttttaaa      480
231      tcaggaaatc gacctcatct tctctctcct cgctctcttc ccctgaaccc ccagtccgca      540
232      tgcactcaca ctctttggcc ttttccctca gtcccgggct cctcttttgt aaatagattt      600
233      gtaggtgtct aagtcacgtc ccaccctcac tcttcccgag gagaggagac agggctagga      660
234      tcccaccgca ccgcgggcca taaacacttg gctgcggcgg ccgc      704
236 <210> SEQ ID NO: 19
237 <211> LENGTH: 263
238 <212> TYPE: DNA
239 <213> ORGANISM: Homo sapiens
240 <400> SEQUENCE: 19
241      gaggaagatg aggtcgattt cctgatttaa aaaatcgtec aagccccgtg gtccagctta      60
242      aggtcctcgg ttacatgcgc cgctcagagc aggtcacttt ctgccttcca cgctctcctt      120
243      caaggaagcc ccatgtgggt agctttcaat atcgaggtt cttactctc tgcctctata      180
244      agtcaaacc caccaacgat cgggcaagta aacccccctc ctgcgcgact tcggaactgg      240
245      cgagagttca gcgcagatgg gcc      263
247 <210> SEQ ID NO: 20
248 <211> LENGTH: 203
249 <212> TYPE: DNA
250 <213> ORGANISM: Homo sapiens
251 <220> FEATURE:
252 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
253      position 16
254 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
255      position 78
256 <400> SEQUENCE: 20
W--> 257      gaggagcccg ggactnaggg aaaaggccaa agagtgtgag tgcagtcgga ctgggggttc      60
W--> 258      aggggaagag gacgagngg aggaagatga ggtcgatttc ctgatttaaa aaatcgccca      120
259      agccccgtg tccagcttaa ggtcctcggg tacatgcgcc gctcagagca ggtcactttc      180
260      tgccttccac gtcctccttc aag      203
262 <210> SEQ ID NO: 21

```

RAW SEQUENCE LISTING ERROR SUMMARY

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Input Set : N:\Crf3\RULE60\10082659.raw

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 2,96

Seq#:3; N Pos. 71,73,76,81,93,138,187,191,192,207,212,223

Seq#:4; N Pos. 2

Seq#:20; N Pos. 16,78

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 31,32,33,34

Seq#:2; Line(s) 41,42,43,44,45

Seq#:3; Line(s) 75,76,77,78

Seq#:4; Line(s) 88,89,90,91,92,93

Seq#:5; Line(s) 100,101

Seq#:6; Line(s) 108,109

Seq#:7; Line(s) 116

Seq#:8; Line(s) 123

Seq#:17; Line(s) 212,213,214,215,216

Seq#:18; Line(s) 223,224,225,226,227,228,229,230,231,232,233,234

Seq#:19; Line(s) 241,242,243,244,245

Seq#:20; Line(s) 257,258,259,260

Seq#:21; Line(s) 267,268,269

VARIABLE LOCATION SUMMARY

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Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.

Use of <220> to <223> is MANDATORY if n's or Xaa's are present.

in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:1; N Pos. 2,96

Seq#:3; N Pos. 71,73,76,81,93,138,187,191,192,207,212,223

Seq#:4; N Pos. 2

Seq#:20; N Pos. 16,78

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/082,659

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Input Set : N:\Crf3\RULE60\10082659.raw

Output Set: N:\CRF3\03132002\J082659.raw

L:31 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:1
L:31 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:1
L:31 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:32 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:1
L:32 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:1
L:32 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:60
L:76 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:3
L:76 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:3
L:76 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:60
L:77 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:3
L:77 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:3
L:77 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:120
L:78 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:3
L:78 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:3
L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:180
L:88 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:4
L:88 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:4
L:88 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:257 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:20
L:257 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:20
L:257 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:258 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:20
L:258 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:20
L:258 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:60